

APPROVED	O.G. FIG.
BY	CLASS SUBCLASS
DRAFTSMAN	

Flag-peptide GAD65 Sgf I IA2 Sgf I PPINS poly-his
 DYKDDDDK-----KKKRRPRKKK-----KKKRRPRKKK-----CNGSHHHHHH

FIG. 1a

Flag-peptide GAD65 Not I IA2 Not I PPINS poly-his
 DYKDDDDK-----KKKRRSRKKK-----KKKRRSRKKK-----CNGSHHHHHH

FIG. 1b

APPROVED	O.G. FIG.
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1A2 Underlined aa 771-979 Accession No. L18983

MRRPRPGGLGGSGGLRLLCLLLSSRPGGCSAVSAHGCLFDRRLCSHLEVCIQDGLFGCQVGVGQARPLLQVTSPVLQRL
 QGVLRQLMSQGLSWHDDLTYVISQEMERIPRLRPEPRPRDRSLAPKRPGPAGELLQDIPTGSAPAAQHRLPQPPVGKGG
 AGASSLSPLQAEALLPLEHLLPPQPPHPSLSYEPALLQPYLFHQFGRDGSRVSEGSPGMVSVGPLPKAEAPALFSRTASKGI
 FGDHPGHSYGDLPGPSPAQLFQDSGLLYLAQELPAPSRARVPRLPEQSSSRAEDSPEGYEKEGLDGRGEPKASPAVQPDAAAL
 QRLAAVLAGYGVVELRQLTPEQLSTLLTLLQLLPGAGRNPGGVNVGADIKKTMEGPVEGRDTAELPARTSPMPGHPTASPT
 SSEVQQVPSVSSEPPKAA RPPVTPVLLLEKKSPLGQSQTVAGQPSARPAEEYGYIVTDQKPLSLAAGVKLLEILAEHVHMSS
 GSFINISVVGPAITFRIRHNEQNLSLADVTQAGLVKSELAQTGLQILQTGVGQREEAAVLPQTAHSTSPMRSVLLTLVALA
 GVA GLLVALAVALCVRQHARQQDKERLAALGPGEAGHDITFEYQDLQRQHMAATKSLFNRAEGPEPSRVSSVSSQFSDAAQ
 ASPSSHSTPSWCEPAQANMDISTGHMILAYMEDHLNRDRLAKWQALCAYQAEPNTCATAQEGENIKNRHPDPLPYDH
 ARIKLKVESPSRSDYINASPIEHDPMPAYIATOGPLSHITADFQWQMVWESGCTVIVMLTPLVEDGVKQCDRYWPDEGASLY
 HVYEVNLYSEHIWCEDEFVRSFYLNKVNQTOETRTLTOFHFLSWPAEGTPASTRPLLDFFRRKVNKCVRGRSCPIIVHCSDGAGR
 TGTYLIDMVLNRMAGVKKEIDIAATLEHVRDORPGLYRSKDQFEFALTAVAEVNAILKALPO

FIG. 2a

GAD65 Underlined aa102-585 Accession No. M74826

MASPGSFWFSGSEDGSDSENPGTARAWCQVAQKFTGGIGNKLCALLYGDAEKPAESGGSQPPRAAARKAACACDQKPCS
 CSKVDVNYAFLHATDLLPACDGERPTLAFLODMVNILLOYYVVKSEDRSTKVIDFHYPNELLQEYNWELADQPONLEILMHC
 QTTLKYAIKTGHPRYFENQLSTGLDMVGLAADWLTTSTANTNMFTYEIAPVFVLLLEYVTLKKMREIIGWPGSGDGFSPGGAIS
 NMYAMMIARFKMFPEVKEKGMAALPRLJAFTSEHSHFSLKKGAAALGIGTDSVILKCDERGMIPSDLERRILEAKOKGFVPP
 LVSATAGTTVYGAFDPLLA VADICKKYKIWMHVDAAWGGGLMSRKHKKWKLSGVERANSVTWNPHKMMGVPLQCSALLY
 REEGLMONCNQMHASYLEQQDKHYDLSYDTGDKALQCGRHVDVFKLWLMWRAKGTTFEAVHVDKCLEAEYLYNIKNR
 EGYEMVFDGKPOHTNVCFWYIPPSLRTLEDNEERMRLSKVAPVIKARMMEYGTITMVSYOPLGDKVNFRRMVISNPAAATHQ
 DIDFLIEEIERLGODL

FIG. 2b

Translation Human preproinsulin.
 EMBL accession nr. v00565

MALWMRLLPLLALLALWGPDPAAAFVNQHLCGSHLVEALYLVCGERGFFYT
 PKTRREAEDLQVGQVELGGGPGAGSLQPLALEGSLQKRGIVEQCCTSICSLYQ
 LENYCN

FIG. 2c

Human GAD65 nucleotide sequence

M74826 Length: 2457 September 1, 1995 12:22 Type: N Check: 8038 ..

APPROVED	O.G.FIG.
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36270" 66EST060

1 ACCCGCCCTC GCCGCTCGGC CCCGCGCGTC CCCGCGCGTG CCCTCCTCCC
51 GCCACACGGC ACGCACGCGC GCGCAGGGCC AAGCCGAGGC AGCCGCCCCG
101 AGCTCGCACT CGCTGGCGAC CTGCTCCAGT CTCCAAAGCC GATGGCATCT
151 CCGGGCTCTG GCTTTTGGTC TTTCGGGTCG GAAGATGGCT CTGGGGATTC
201 CGAGAATCCC GGCACAGCGC GAGCCTGGTG CCAAGTGGCT CAGAAGTTCA
251 CGGGCGGCAT CGGAAACAAA CTGTGCGCCC TGCTCTACGG AGACGCCGAG
301 AAGCCGGCGG AGAGCGGCGG GAGCCAACCC CCGCGGGCCG CCGCCCCGAA
351 GGCCGCCTGC GCCTGCGACC AGAAGCCCTG CAGCTGCTCC AAAGTGGATG
401 TCAACTACGC GTTCTCCAT GCAACAGACC TGCTGCCGGC GTGTGATGGA

451 GAAAGGCCCA CTTTGGCGTT TCTGCAAGAT GTTATGAACA TTTACTTCA
501 GTATGTGGTG AAAAGTTTCG ATAGATCAAC CAAAGTGATT GATTTCATT
551 ATCCTAATGA GCTTCTCCAA GAATATAATT GGAATTGGC AGACCAACCA
601 CAAAATTTGG AGGAAATTTT GATGCATTGC CAAACAACCT TAAAATATGC
651 AATTAAAACA GGGCATCCTA GATACTTCAA TCAACTTTCT ACTGGTTTGG
701 ATATGGTTGG ATTAGCAGCA GACTGGCTGA CATCAACAGC AAATACTAAC
751 ATGTTACCT ATGAAATTGC TCCAGTATTT GTGCTTTTGG AATATGTCAC
801 ACTAAAGAAA ATGAGAGAAA TCATTGGCTG GCCAGGGGGC TCTGGCGATG
851 GGATATTTTC TCCCGGTGGC GCCATATCTA ACATGTATGC CATGATGATC
901 GCACGCTTTA AGATGTTCCC AGAAGTCAAG GAGAAAGGAA TGGCTGCTCT
951 TCCCAGGCTC ATTGCCTTCA CGTCTGAACA TAGTCATTTT TCTCTAAGA
1001 AGGGAGCTGC AGCCTTAGGG ATTGGAACAG ACAGCGTGAT TCTGATTAAA
1051 TGTGATGAGA GAGGGAAAAT GATTCCATCT GATCTTGAAA GAAGGATTCT
1101 TGAAGCCAAA CAGAAAGGGT TTGTTCCCTT CCTCGTGAGT GCCACAGCTG
1151 GAACCACCGT GTACGGAGCA TTTGACCCCC TCTTAGCTGT CGCTGACATT
1201 TGCAAAAAGT ATAAGATCTG GATGCATGTG GATGCAGCTT GGGGTGGGG
1251 ATTACTGATG TCCCGAAAAC ACAAGTGGAA ACTGAGTGGC GTGGAGAGGG

FIG. 3a

APPROVED BY DRAFTSMAN	O.G. FIG.
	CLASS / SUBCLASS

8662FD-65EST060

1301 CCAACTCTGT GACGTGGAAT CCACACAAGA TGATGGGAGT CCCTTTGCAG
 1351 TGCTCTGCTC TCCTGGTTAG AGAAGAGGGA TTGATGCAGA ATTGCAACCA
 1401 AATGCATGCC TCCTACCTCT TTCAGCAAGA TAAACATTAT GACCTGTCTT
 1451 ATGACACTGG AGACAAGGCC TTACAGTGCG GACGCCACGT TGATGTTTTT
 1501 AAACATATGGC TGATGTGGAG GGCAAAGGGG ACTACCGGGT TTGAAGCGCA
 1551 TGTTGATAAA TGTTTGGAGT TGGCAGAGTA TTTATACAAC ATCATAAAAA
 1601 ACCGAGAAGG ATATGAGATG GTGTTTGATG GGAAGCCTCA GCACACAAAT
 1651 GTCTGCTTCT GGTACATTCC TCCAAGCTTG CGTACTCTGG AAGACAATGA
 1701 AGAGAGAATG AGTCGCCTCT CGAAGGTGGC TCCAGTGATT AAAGCCAGAA
 1751 TGATGGAGTA TGGAACCACA ATGGTCAGCT ACCAACCCTT GGGAGACAAG

 1801 GTCAATTTCT TCCGCATGGT CATCTCAAAC CCAGCGGCAA CTCACCAAGA
 1851 CATTGACTTC CTGATTGAAG AAATAGAACG CCTTGGACAA GATTTATAAT
 1901 AACCTTGCTC ACCAAGCTGT TCCACTTCTC TAGAGAACAT GCCCTCAGCT
 1951 AAGCCCCCTA CTGAGAAACT TCCTTTGAGA ATTGTCGAC TTCACAAAAT
 2001 GCAAGGTGAA CACCACTTTG TCTCTGAGAA CAGACGTTAC CAATTATGGA
 2051 GTGTCACCAG CTGCCAAAAT CGTAGGTGTT GGCTCTGCTG GTCACTGGAG
 2101 TAGTTGCTAC TCTTCAGAAT ATGGACAAAG AAGGCACAGG TGTAATATA
 2151 GTAGCAGGAT GAGGAACCTC AAACCTGGGT TCATTTGCAC GTGCTCTTCT
 2201 GTTCTCAAAT GCTAAATGCA AACACTGTGT ATTTATTAGT TAGGTGTGCC
 2251 AAACCTACCGT TCCCAAATTG GTGTTTCTGA ATGACATCAA CATTCCCCCA
 2301 ACATTACTCC ATTACTAAAG ACAGAAAAAA ATAAAAACAT AAAATATACA
 2351 AACATGTGGC AACCTGTTCT TCCTACCAAA TATAAACTTG TGTATGATCC
 2401 AAGTATTTTA TCTGTGTTGT CTCTCTAAAC CCAAATAAAT GTGTAAATGT
 2451 GGACACA

FIG. 3b

Human IA-2 nucleotide sequence

L18983 Length: 3613 November 20, 1997 16:45 Type: N Check: 6409 ..

APPROVED	O.G.FIG.
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BY	
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1 CAGCCCCTCT GGCAGGCTCC CGCCAGCGTC GCTGCGGCTC CGGCCCCGGA
51 GCGAGCGCCC GGAGCTCGGA AAGATGCGGC GCCCGCGGCG GCCTGGGGGT
101 CTCGGGGGAT CCGGGGGTCT CCGGCTGCTC CTCTGCCTCC TGCTGCTGAG
151 CAGCCGCCCC GGGGGCTGCA GCGCCGTTAG TGCCACGGC TGTCTATTG
201 ACCGCAGGCT CTGCTCTCAC CTGGAAGTCT GTATTACAGGA TGGCTTGTTT
251 GGGCAGTGCC AGGTGGGAGT GGGGCAGGCC CGGCCCTTT TGCAAGTCAC
301 CTCCCCAGTT CTCCAACGCT TACAAGGTGT GCTCCGACAA CTCATGTCCC
351 AAGGATTGTC CTGGCACGAT GACCTCACCC AGTATGTGAT CTCTCAGGAG
401 ATGGAGCGCA TCCCCAGGCT TCGCCCCCA GAGCCCCGTC CAAGGGACAG
451 GTCTGGCTTG GCACCAAGA GACCTGGTCC TGCTGGAGAG CTGCTTTTAC
501 AGGACATCCC CACTGGCTCC GCCCTGCTG CCCAGCATCG GCTTCCACAA
551 CCACCAAGTG GCAAAGGTGG AGCTGGGGCC AGCTCCTCTC TGTCCTCTT
601 GCAGGCTGAG CTGCTCCCGC CTCTCTTGA GCACCTGCTG CTGCCCCAC
651 AGCTCCCCA CCCTTCACTG AGTTACGAAC CTGCCTTGCT GCAGCCCTAC
701 CTGTTCCACC AGTTTGGCTC CCGTGATGGC TCCAGGGTCT CAGAGGGCTC
751 CCCAGGGATG GTCAGTGTG GCCCCCTGCC CAAGGCTGAA GCCCCTGCCC
801 TCTTCAGCAG AACTGCCTCC AAGGGCATAT TTGGGGACCA CCCTGGCCAC
851 TCCTACGGGG ACCTTCAGG GCCTTCACCT GCCCAGCTTT TTCAAGACTC
901 TGGGCTGCTC TATCTGGCCC AGGAGTTGCC AGCACCCAGC AGGGCCAGGG
951 TGCCAAGGCT GCCAGAGCAA GGGAGCAGCA GCCGGGCAGA GGAATCCCCA
1001 GAGGGCTATG AGAAGGAAGG ACTAGGGGAT CGTGGAGAGA AGCCTGCTTC
1051 CCCAGCTGTG CAGCCAGATG CGGCTCTGCA GAGGCTGGCC GCTGTGCTGG
1101 CGGGCTATGG GGTAGAGCTG CGTCAGCTGA CCCCTGAGCA GCTCTCCACA
1151 CTCCTGACCC TGCTGCAGCT ACTGCCCAAG GGTGCAGGAA GAAATCCGGG
1201 AGGGGTTGTA AATGTTGGAG CTGATATCAA GAAAACAATG GAGGGGCCGG
1251 TGGAGGGCAG AGACACAGCA GAGCTTCCAG CCCGCACATC CCCCATGCCT

FIG. 3c

APPROVED	O.G. FIG.
	CLASS SUBCLASS
BY	
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1301 GGACACCCCA CTGCCAGCCC TACCTCCAGT GAAGTCCAGC AGGTGCCAAG
 1351 CCCTGTCTCC TCTGAGCCTC CCAAAGCTGC CAGACCCCCT GTGACACCTG
 1401 TCCTGCTAGA GAAGAAAAGC CCACTGGGCC AGAGCCAGCC CACGGTGGCA
 1451 GGACAGCCCT CAGCCCCGCC AGCAGCAGAG GAATATGGCT ACATCGTCAC
 1501 TGATCAGAAG CCCCTGAGCC TGGCTGCAGG AGTGAAGCTG CTGGAGATCC
 1551 TGGCTGAGCA TGTGCACATG TCCTCAGGCA GCTTCATCAA CATCAGTGTG
 1601 GTGGGACCAG CCCTCACCTT CCGCATCCGG CACAATGAGC AGAACCTGTC
 1651 TTTGGCTGAT GTGACCCAAC AAGCAGGGCT GGTGAAGTCT GAACTGGAAG
 1701 CACAGACAGG GCTCCAAATC TTGCAGACAG GAGTGGGACA GAGGGAGGAG
 1751 GCAGCTGCAG TCCTTCCCCA AACTGCGCAC AGCACCTCAC CCATGCGCTC
 1801 AGTGCTGCTC ACTCTGGTGG CCCTGGCAGG TGTGGCTGGG CTGCTGGTGG
 1851 CTCTGGCTGT GGCTCTGTGT GTGCGGCAGC ATGCGCGGCA GCAAGACAAG
 1901 GAGCGCCTGG CAGCCCTGGG GCCTGAGGGG GCCCATGGTG ACACTACCTT
 1951 TGAGTACCAG GACCTGTGCC GCCAGCACAT GGCCACGAAG TCCTTGTTC
 2001 ACCGGGCAGA GGGTCCACCG GAGCCTTCAC GGGTGAGCAG TGTGTCCTCC
 2051 CAGTTCAGCG ACGCAGCCCA GGCCAGCCCC AGCTCCCACA GCAGACCCCC
 2101 GTCCTGGTGC GAGGAGCCGG CCAAGCCAA CATGGACATC TCCACGGGAC
 2151 ACATGATTCT GGCATACATG GAGGATCACC TCGGAACCG GGACCGCCTT
 2201 GCCAAGGAGT GGCAGGCCCT CTGTGCCTAC CAAGCAGAGC CAAACACCTG
 2251 TGCCACCGCG CAGGGGGAGG GCAACATCAA AAAGAACCGG CATCCTGACT
 2301 TCCTGCCCTA TGACCATGCC CGCATAAAAC TGAAGGTGGA GAGCAGCCCT
 2351 TCTCGGAGCG ATTACATCAA CGCCAGCCCC ATTATTGAGC ATGACCCTCG
 2401 GATGCCAGCC TACATAGCCA CGCAGGGCCC GCTGTCCCAT ACCATCGCAG
 2451 ACTTCTGGCA GATGGTGTGG GAGAGCGGCT GCACCGTCAT CGTCATGCTG
 2501 ACCCCGCTGG TGGAGGATGG TGTCAAGCAG TGTGACCGCT ACTGGCCAGA
 2551 TGAGGGTGCC TCCCTCTACC ACGTATATGA GGTGAACCTG GTGTGCGGAGC
 2601 ACATCTGGTG CGAGGACTTT CTGGTGCGGA GCTTCTACCT GAAGAACGTG
 2651 CAGACCCAGG AGACGCGCAC GCTCACGCAG TTCCAATTCC TCAGCTGGCC

FIG. 3d

APPROVED BY DRAFTSMAN	OG. FIG.
	CLASS SUBCLASS

86270" 66EST060

2701 GGCAGAGGGC ACACCGGCCT CCACGCGGCC CCTGCTGGAC TTCCGCAGGA
 2751 AGGTGAACAA GTGCTACCGG GGCCGCTCCT GCCCCATCAT CGTGCACTGC
 2801 AGTGATGGTG CGGGGAGGAC CGGCACCTAC ATCCTCATCG ACATGGTCCT
 2851 GAACCGCATG GCAAAAGGAG TGAAGGAGAT TGACATCGCT GCCACCCTGG
 2901 AGCATGTCCG TGACCAGCGG CCTGGCCTTG TCCGCTCTAA GGACCAGTTT
 2951 GAATTTGCCC TGACAGCCGT GCGGAGGAA GTGAATGCCA TCCTCAAGGC
 3001 CCTGCCCCAG TGAGACCCTG GGGCCCCTTG GCGGGCAGCC CAGCCTCTGT
 3051 CCCTCTTTGC CTGTGTGAGC ATCTCTGTGT ACCCACTCCT CACTGCCCCA
 3101 CCAGCCACCT CTTGGGCATG CTCAGCCCTT CCTAGAAGAG TCAGGAAGGG
 3151 AAAGCCAGAA GGGGCACGCC TGCCAGCCT CGCATGCCAG AGCCTGGGGC
 3201 ATCCCAGAGC CCAGGGCATC CCATGGGGGT GCTGCAGCCA GGAGGAGAGG
 3251 AAAGGACATG GGTAGCAATT CTACCCAGAG CTTTCTCCTG CCTACATTCC
 3301 CTGGCCTGGC TCTCCTGTAG CTCTCCTGGG GTTCTGGGAG TTCCTGAAC
 3351 ATCTGTGTGT GTCCCCCTAT GCTCCAGTAT GGAAGAATGG GGTGGAGGGT
 3401 CGCCACACCC GGCTCCCCCT GCTTCTCAGC CCCGGGCCTG CCTCTGACTC
 3451 AACTTGGGC GCTCTGCCCT CCCTGGCCTC ACGCCAGCC TGGTCCCACC
 3501 ACCCTCCCAC CATGCGCTGC TCAACCTCTC TCCTTCTGGC GCAAGAGAAC
 3551 ATTTCTAGAA AAAACTACTT TTGTACCAGT GTGAATAAAG TTAGTGTGTT
 3601 GTCTGTGCAG CTG

FIG. 3e

PREPROINSULINI

Exon sequences, i.e. sequences to be used in the patent are underlined and represent exon sequences.

V00565 Length: 4992 December 18, 1997 17:50 Type: N Check: 9721 ..

APPROVED	O.G.FIG.
	CLASS SUBCLASS
BY	
DRAFTSMAN	

365210" 66EST060

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1  CTCGAGGGGC CTAGACATTG CCCTCCAGAG AGAGCACCCA ACACCCTCCA
51  GGCTTGACCG GCCAGGGTGT CCCCTTCCTA CCTTGGAGAG AGCAGCCCCA
101 GGGCATCCTG CAGGGGGTGC TGGGACACCA GCTGGCCTTC AAGGTCTCTG
151 CCTCCCTCCA GCCACCCAC TACACGCTGC TGGGATCCTG GATCTCAGCT
201 CCCTGGCCGA CAACACTGGC AAACCTCCTAC TCATCCACGA AGGCCCTCCT
251 GGGCATGGTG GTCCTTCCCA GCCTGGCAGT CTGTTCTCTCA CACACCTTGT
301 TAGTGCCAG CCCCTGAGGT TGCAGCTGGG GGTGTCTCTG AAGGGCTGTG
351 AGCCCCCAGG AAGCCCTGGG GAAGTGCCTG CTTGCCTCC CCCCggccct
401 GCCAGCGCCT GGCTCTGCCC TCCTACCTGG GCTCCCCCA TCCAGCCTCC
451 CTCCCTACAC ACTCCTCTCA AGGAGGCACC CATGTCCTCT CCAGCTGCCG
501 GGCCTCAGAG CACTGTGGCG TCCTGGGGCA GCCACCGCAT GTCCTGCTGT
551 GGCATGGCTC AGGGTGGAAA GGGCGGAAGG GAGGGGTCCT GCAGATAGCT
601 GGTGCCCCACT ACCAAACCCG CTCGGGGCAG GAGAGCCAAA GGCTGGGTGT
651 GTGCAGAGCG GCCCCGAGAG GTTCCGAGGC TGAGGCCAGG GTGGGACATA
701 GGGATGCGAG GGGCCGGGGC ACAGGATACT CCAACCTGCC TGCCCCCATG
751 GTCTCATCCT CCTGCTTCTG GGACCTCCTG ATCCTGCCCC TGGTGCTAAG
801 AGGCAGGTAA GGGGCTGCAG GCAGCAGGGC TCGGAGCCCA TGCCCCCTCA
851 CCATGGGTCA GGCTGGACCT CCAGGTGCCT GTTCTGGGGA GCTGGGAGGG
901 CCGGAGGGGT GTACCCAGG GGCTCAGCCC AGATGACACT ATGGGGGTGA
951 TGGTGTGATG GGACCTGGCC AGGAGAGGGG AGATGGGCTC CCAGAAGAGG
1001 AGTGGGGGCT GAGAGGGTGC CTGGGGGGCC AGGACGGAGC TGGGCCAGTG
1051 CACAGCTTCC CACACCTGCC CACCCCCAGA GTCCTGCCGC CACCCCCAGA
1101 TCACACGGAA GATGAGGTCC GAGTGGCCTG CTGAGGACTT GCTGCTTGTC
1151 CCCAGGTCCC CAGGTGATGC CCTCCTTCTG CCACCCTGGG GAGCTGAGGG
1201 CCTCAGCTGG GGCTGCTGTC CTAAGGCAGG GTGGGAAC TA GGCAGCCAGC
1251 AGGGAGGGGA CCCCTCCCTC ACTCCCACTC TCCACCCCC ACCACCTTGG
1301 CCCATCCATG GCGGCATCTT GGGCCATCCG GGAAGTGGGA CAGGGGTCTT
1351 GGGGACAGGG GTCCGGGGAC AGGGTCCTGG GGACAGGGGT GTGGGGACAG
  
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FIG. 3f

APPROVED	O.G.FIG.
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DRAFTSMAN	

855210" 66E3F066

1401 GGGTCTGGGG ACAGGGGTGT GGGGACAGGG GTGTGGGGAC AGGGGTCTGG
 1451 GGACAGGGGT GTGGGGACAG GGGTCCGGGG ACAGGGGTGT GGGGACAGGG
 1501 GTCTGGGGAC AGGGGTGTGG GGACAGGGGT GTGGGGACAG GGGTCTGGGG
 1551 ACAGGGGTGT GGGGACAGGG GTCCTGGGGA CAGGGGTGTG GGGACAGGGG
 1601 TGTGGGGACA GGGGTGTGGG GACAGGGGTG TGGGGACAGG GGTCTCTGGG
 1651 ATAGGGGTGT GGGGACAGGG GTGTGGGGAC AGGGGTCCCG GGGACAGGGG
 1701 TGTGGGGACA GGGGTGTGGG GACAGGGGTC CTGGGGACAG GGGTCTGAGG
 1751 ACAGGGGTGT GGGCACAGGG GTCCTGGGGA CAGGGGTCCT GGGGACAGGG
 1801 GTCCTGGGGA CAGGGGTCTG GGGACAGCAG CGCAAAGAGC CCCGCCCTGC
 1851 AGCCTCCAGC TCTCTGGTC TAATGTGGAA AGTGGCCCAG GTGAGGGCTT
 1901 TGCTCTCCTG GAGACATTTG CCCCCAGCTG TGAGCAGGGA CAGGTCTGGC
 1951 CACCGGGCCC CTGGTTAAGA CTCTAATGAC CCGCTGGTCC TGAGGAAGAG
 2001 GTGCTGACGA CCAAGGAGAT CTTCCACAG ACCCAGCACC AGGGAAATGG
 2051 TCCGGAAATT GCAGCCTCAG CCCCCAGCCA TCTGCCGACC CCCCCACCC
 2101 GCCCTAATGG GCCAGGCGGC AGGGGTTGAC AGGTAGGGGA GATGGGCTCT
 2151 GAGACTATAA AGCCAGCGGG GGCCCAGCAG CCTCAGCCC TCCAGGACAG
 2201 GCTGCATCAG AAGAGGCCAT CAAGCAGGTC TGTTC AAGG GCCTTTGCGT
 2251 CAGGTGGGCT CAGGGTTCCA GGGTGGCTGG ACCCCAGGCC CCAGCTCTGC
 2301 AGCAGGGAGG ACGTGGCTGG GCTCGTGAAG CATGTGGGGG TGAGCCCAGG
 2351 GGCCCCAAGG CAGGGCACCT GGCCTTCAGC CTGCCTCAGC CCTGCCTGTC
 2401 TCCCAGATCA CTGTCCTTCT GCCATGGCCC TGTGGATGCG CCTCCTGCCC
 2451 CTGCTGGCGC TGCTGGCCCT CTGGGGACCT GACCCAGCCG CAGCCTTTGT
 2501 GAACCAACAC CTGTGCGGCT CACACCTGGT GGAAGCTCTC TACCTAGTGT
 2551 GCGGGGAACG AGGCTTCTTC TACACACCCA AGACCCGCCG GGAGGCAGAG
 2601 GACCTGCAGG GTGAGCCAAC CGCCCATTGC TGCCCCTGGC CGCCCCCAGC
 2651 CACCCCCTGC TCCTGGCGCT CCCACCCAGC ATGGGCAGAA GGGGGCAGGA
 2701 GGCTGCCACC CAGCAGGGGG TCAGGTGCAC TTTTTTAAAA AGAAGTTCTC
 2751 TTGGTCACGT CCTAAAAGTG ACCAGCTCCC TGTGGCCCAG TCAGAATCTC
 2801 AGCCTGAGGA CGGTGTTGGC TTCGGCAGCC CCGAGATACA TCAGAGGGTG
 2851 GGCACGCTCC TCCCTCCACT CGCCCCTCAA ACAAATGCCC CGCAGCCCAT

FIG. 3g

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FIG. 3h

APPROVED	O.G. FIG.
	CLASS SUBCLASS
BY	DRAFTSMAN

4401 CTTGCGGCCC TTAGCCCACC CCCTCCCAGT AAGCAGGGGC TGCTTGGCTA
4451 GGCTTCCTTT TGCTACAGAC CTGCTGCTCA CCCAGAGGCC CACGGGCCCT
4501 AGTGACAAGG TCGTTGTGGC TCCAGGTCCT TGGGGGTCCT GACACAGAGC
4551 CTCTTCTGCA GCACCCCTGA GGACAGGGTG CTCCGCTGGG CACCCAGCCT
4601 AGTGGGCAGA CGAGAACCTA GGGGCTGCCT GGGCCTACTG TGGCCTGGGA
4651 GGTGAGCGGG TGACCCTAGC TACCCTGTGG CTGGGCCAGT CTGCCTGCCA
4701 CCCAGGCCAA ACCAATCTGC ACCTTTCCTG AGAGCTCCAC CCAGGGCTGG
4751 GCTGGGGATG GCTGGGCCTG GGGCTGGCAT GGGCTGTGGC TGCAGACCAC
4801 TGCCAGCTTG GGCCTCGAGG CCAGGAGCTC ACCCTCCAGC TGCCCCGCCT
4851 CCAGAGTGGG GGCCAGGGCT GGGCAGGCGG GTGGACGGCC GGACACTGGC
4901 CCCGGAAGAG GAGGGAGGCG GTGGCTGGGA TCGGCAGCAG CCGTCCATGG
4951 GAACACCCAG CCGGCCCCAC TCGCACGGGT AGAGACAGGC GC

FIG. 3i

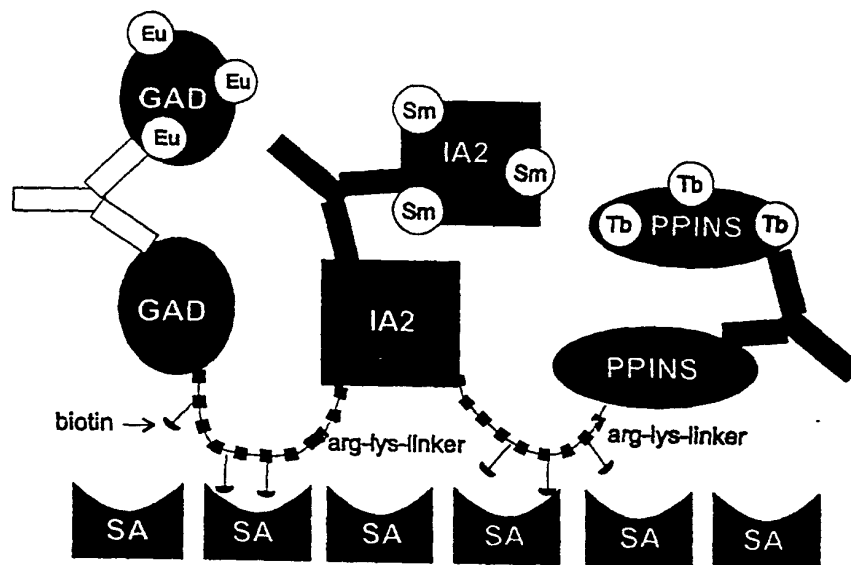


FIG. 4

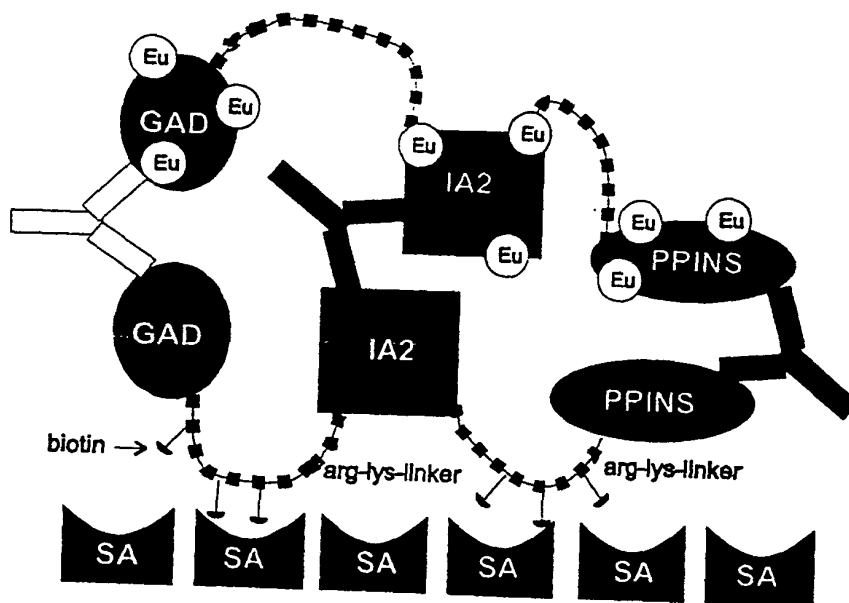


FIG. 5